SEQUENCE LISTING

<110>	INSTITUT DE LA RECHERCHE POUR LE DEVELOPPEMENT (IRD)	
<120>	NOVEL AGENTS FOR THE PREVENTION OF LEISHMANIOSIS	
<130>	1721-116	
<140>	10/579,749	
<141>	2006-05-18	
<150>	PCT/FR04/02955	
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<150>	FR 03 13 555	
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<151>	2004-06-25	
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Val Val Ala Leu Leu Cys Thr Ser Ser Ala Pro Val Ala Arg Ala 20 25 30

Ala Gly Thr Ser Asp Phe Thr Glu Ala Gln Gln Thr Asn Thr Leu Thr 35 40 45

Val Leu Gln Ala Phe Ala Arg Ala Ile Pro Ala Leu Gly Asp Thr Trp 50 55 60

Thr Gly Ser Asp Phe Cys Ser Trp Lys His Ile Ile Cys Asp Ser Pro

80

Gly Val Gly Val Trp Met Gly Asp Val Asp Tyr Thr Gly Thr Leu Pro 85 90 95

70

Glu Met Pro Ala Ser Val Asp Tyr Lys Asp Val Met Ile Thr Glu Leu 100 105 110

Asn Phe Ser Ala Met Gly Gln Gly Leu Ser Gly Thr Leu Pro Pro Ser 115 120 125

Trp Ser Ser Leu Thr Ser Leu Ile Ser Leu Cys Ile Glu Lys Ser Glu 130 135 140

Lys Val Thr Gly Thr Leu Pro Ala Gln Trp Ser Ser Met Thr Ser Leu 145 150 155 160

Asp Asn Leu Asn Leu His Asp Thr Ala Val Ser Gly Thr Leu Pro Ala 165 170 175

Gln Trp Ser Ser Met Lys Gln Leu Thr Val Leu Asp Leu Glu Gly Thr 180 185 190

Lys Val Ser Gly Thr Leu Pro Ser Glu Trp Ser Gly Met Ala Lys Ala. 195 200 205

Glu Ala Val Gln Leu Glu Asn Cys Gly Leu Ser Gly Ser Leu Pro Pro 210 215 220

Ser Trp Ser Ala Met Pro Lys Leu Arg Ile Val Ser Leu Ser Gly Asn 225 230 235 240

His Phe Cys Gly Cys Val Pro Asp Ser Trp Arg Glu Lys Asp Arg Leu 245 250 255

Asp Val Thr Ile Glu Glu Trp His Met Gly Glu Asp Cys Lys Leu Ala 260 265 270

Asn Ala Cys Arg Pro Thr Ala Ala Pro Gly Thr Thr Thr Thr Asn Pro 275 280 285

Pro Thr Thr Thr Gly Thr Pro Ala Ala Ser Ser Thr Pro Ser Pro Gly 290 295 300

Ser Gly Cys Glu Val Asp Gly Cys Glu Val Cys Glu Gly Asp Ser Ala 305 310 315 320 Ala Arg Cys Ala Arg Cys Arg Glu Gly Tyr Ser Leu Thr Asp Glu Lys 325 330 335

Thr Cys Leu Ala Asn His Asp Gly Gly Val Ala Ala Ala Ser Ser Gly 340 345 350

Ala Val Ala Ala Ala Val Trp Ala Ala Val Leu Leu Ser Val Gly 355 360 365

Leu Val Ala 370

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Val Asp Tyr Lys Asp Val Met Ile Thr Glu Leu Asn Phe Ser Ala Met 20 25 30

Gly Gln Gly Leu Ser Gly Thr Leu Pro Pro Ser Trp Ser Ser Leu Thr 35 40 45

Ser Leu Ile Ser Leu Cys Ile Glu Lys Ser Glu Lys Val Thr Gly Thr 50 55 60

Leu Pro Ala Gln Trp Ser Ser Met Thr Ser Leu Asp Asn Leu Asn Leu 65 70 75 80

His Asp Thr Ala Val Ser Gly Thr Leu Pro Ala Gln Trp Ser Ser Met 85 90 95

Lys Gln Leu Thr Val Leu Asp Leu Glu Gly Thr Lys Val Ser Gly Thr 100 105 110

Leu Pro Ser Glu Trp Ser Gly Met Ala Lys Ala Glu Ala Val Gln Leu 115 120 125 Glu Asn Cys Gly Leu Ser Gly Ser Leu Pro Pro Ser Trp Ser Ala Met 130 135 140

Pro Lys Leu Arg Ile Val Ser Leu Ser Gly Asn His Phe Cys Gly Cys 145 150 155 160

Val Pro Asp Ser Trp Arg Glu Lys Asp Arg Leu Asp Val Thr Ile Glu 165 170 175

Glu Trp His Met Gly Glu Asp Cys Lys Leu Ala Asn Ala Cys Arg Pro 180 185 190

Thr Ala Ala Pro Gly Thr Thr Thr Thr Asn Pro Pro Thr Thr Thr Gly
195 200 205

Thr Pro Ala Ala Ser Ser Thr Pro Ser Pro Gly Ser Gly Cys Glu Val 210 215 220

Asp Gly Cys Glu Val Cys Glu Gly Asp Ser Ala Ala Arg Cys Ala Arg 225 230 235 240

Cys Arg Glu Gly Tyr Ser Leu Thr Asp Glu Lys Thr Cys Val Ala Asn 245 250 255

His Asp Gly Gly Val Ala Ala Ala Ser Ser Gly Ala Val Ala Ala Ala 260 265 270

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Gly Gln Gly Leu Ser Gly Thr Leu Pro Pro Ser Trp Ser Ser Leu Thr 35 40 45

Ser Leu Met Ser Leu Trp Ile Glu Lys Ser Glu Lys Val Thr Gly Thr 50 55 60

Leu Pro Thr Gln Trp Ser Ser Met Lys Gln Leu Thr Leu Leu His Leu 65 70 75 80

Lys Gly Thr Lys Val Ser Gly Thr Leu Pro Pro Glu Trp Ser Gly Met 85 90 95

Thr Ser Leu Asp Asp Leu Asn Leu His Asp Thr Ala Val Ser Gly Thr

Leu Pro Ala Gln Trp Ser Ser Met Lys Gln Leu Ile Asp Leu Asp Leu 115 120 125

Glu Gly Thr Lys Val Ser Gly Thr Leu Pro Pro Glu Trp Ser Gly Met 130 135

Ala Lys Ala Glu Ala Leu Gln Leu Lys Tyr Cys Asp Leu Ser Gly Ser 145 150 155 160

Leu Pro Pro Ser Trp Ser Ser Met Gln Lys Leu Arg Ile Val Ser Leu 165 170 175

Ser Gly Asn His Phe Cys Gly Cys Val Pro Asp Ser Trp Arg Glu Lys 180 185 190

Asp Arg Leu Asp Val Thr Ile Glu Glu Trp His Met Gly Glu Asp Cys 195 200 205

Lys Leu Ala Asn Ala Cys Arg Pro Thr Ala Ala Pro Gly Thr Thr Thr 210 215 220

Thr Asn Pro Pro Thr Thr Thr Gly Thr Pro Ala Ala Ser Ser Thr Pro 225 230 235 240

Ser Pro Gly Ser Gly Cys Glu Val Asp Gly Cys Glu Val Cys Glu Gly 245 250 255

Asp Ser Ala Ala Arg Cys Ala Arg Cys Arg Glu Gly Tyr Ser Leu Thr 260 265 270

Asp Glu Lys Thr Cys Leu Ala Asn His Asp Gly Gly Val Ala Ala Ala

275 280 285

Ser Ser Gly Ala Val Ala Ala Ala Val Trp Ala Ala Val Leu Leu 290 295 300

Ser Val Gly Leu Val Ala 305 310

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Phe Ala Ala Arg Leu Arg Cys Gly Cys His Ser Leu Trp Ala Arg Trp 20 25 30

Arg Val Ala Pro Gly Gly Ala Arg Ser Ala Gly Gly Gly Ser Ser Val

Phe Leu Tyr Phe Ser Val Pro Cys Cys Pro Pro Lys Lys Lys Lys 50 55 60

Lys Lys Ile Gly Val Trp Met Gly Asp Val Asp Tyr Thr Gly Thr Leu 65 70 75 80

Pro Glu Met Pro Ala Ser Val Asp Tyr Lys Asp Val Met Ile Thr Glu 85 90 95

Leu Asn Phe Gly Ala Met Gly Gln Gly Leu Ser Gly Thr Leu Pro Pro 100 105 110

Ser Trp Ser Ser Met Lys Gln Leu Ile Asp Leu Asp Leu Glu Gly Thr 115 120 125

Lys Val Ser Gly Thr Leu Pro Pro Glu Trp Ser Gly Met Ala Lys Ala

Glu Ala Leu Gln Leu Lys Tyr Cys Asp Leu Ser Gly Ser Leu Pro Pro 145 150 155 160 Ser Trp Ser Ser Met Gln Lys Leu Arg Ile Val Ser Leu Ser Gly Asn 165 170 175

His Phe Cys Gly Cys Val Pro Asp Ser Trp Arg Glu Lys Asp Arg Leu 180 185 190

Asp Val Thr Ile Glu Glu Trp His Met Gly Glu Asp Cys Lys Leu Ala 195 200 205

Asn Ala Cys Arg Pro Thr Ala Ala Pro Gly Thr Thr Thr Thr Asn Pro 210 215 220

Pro Thr Thr Gly Thr Pro Ala Ala Ser Ser Thr Pro Ser Pro Gly 225 230 235

Ser Gly Cys Glu Val Asp Gly Cys Glu Val Cys Glu Gly Asp Ser Ala 245 250 255

Ala Arg Cys Ala Arg Cys Arg Glu Gly Tyr Ser Leu Thr Asp Glu Lys 260 265 270

Thr Cys Leu Ala Asn His Asp Gly Gly Val Ala Ala Ala Ser Ser Gly 275 280 285

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Leu Val Ala 305

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Gly Gln Gly Leu Ser Gly Thr Leu Pro Pro Ser Trp Ser Ser Leu Thr 35 40 45

Ser Leu Ile Ser Leu Cys Ile Glu Lys Ser Glu Lys Val Thr Gly Thr 50 55 60

Leu Pro Ala Gln Trp Ser Ser Met Thr Ser Leu Asp Asn Leu Asn Leu 65 70 75 80

His Asp Thr Ala Val Ser Gly Thr Leu Pro Pro Glu Trp Ser Gly Met 85 90 95

Leu Pro Ala Gln Trp Ser Ser Met Lys Gln Leu Ile Asp Leu Asp Leu 115 120 125

Glu Gly Thr Lys Val Ser Gly Thr Leu Pro Pro Glu Trp Ser Gly Met 130 140

Ala Lys Ala Glu Ala Leu Gln Leu Lys Tyr Cys Asp Leu Ser Gly Ser 145 150 155 160

Leu Pro Pro Ser Trp Ser Ser Met Gln Lys Leu Arg Ile Val Ser Leu 165 170 175

Ser Gly Asn His Phe Cys Gly Cys Val Pro Asp Ser Trp Arg Glu Lys 180 185 190

Asp Arg Leu Asp Val Thr Ile Glu Glu Trp His Met Gly Glu Asp Cys 195 200 205

Lys Leu Ala Asn Ala Cys Arg Pro Thr Ala Ala Pro Gly Thr Thr Thr 210 215 220

Thr Asn Pro Pro Thr Thr Gly Thr Pro Ala Ala Ser Ser Thr Pro 225 230 235

Ser Pro Gly Ser Gly Cys Glu Val Asp Gly Cys Glu Val Cys Glu Gly 245 250 255

Asp Ser Ala Ala Arg Cys Ala Arg Cys Arg Glu Gly Tyr Ser Leu Thr 260 265 270

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Ala Val Lys Asp Asp Phe Thr Ala Ala Gln Arg Thr Asn Thr Leu Ala 35 40 45

Val Leu Glu Ala Phe Gly Arg Ala Ile Pro Glu Leu Gly Lys Leu Trp 50 55 60

Lys Gly Asp Asp Phe Cys Phe Trp Glu Ser Val Val Val Arg Cys Asp 65 70 75 80

Arg Ser Val Leu Gly Gly Lys Ser Val Arg Arg Ile Pro Ala Arg Cys
85 90 95

Arg Arg Cys Leu Trp Thr Ser Thr Thr Arg Pro Ser Trp Ser Ser Thr

Ser Thr Phe Pro Lys Trp Gly Trp Gly Trp Ala Glu Arg Cys Arg Thr 115 120 125

Ala Gly Ala Gly Cys Arg Asp Trp Pro His Leu Arg Cys Arg Ala Ala 130 135

Ala Trp Ala Val Arg Cys Pro Pro Arg Gly Ala Arg Trp Ser Leu Trp 145 150 155 160

Cys Arg Cys Gly Leu Arg Val Val Lys Val Leu Pro Ala Ser Cys Arg 165 170 175

- Leu Ser Gly Ala Arg Trp Asn Arg Trp Glu Ile Ser Ile Cys Met Ala 180 185 190
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- Trp Pro Phe Ser Ile Cys Arg Thr Leu Arg Leu Pro Ala Val Cys Arg 210 215 220
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- Trp Ala Phe Ser Val Trp Arg Val Leu Ser Ser Pro Ala Arg Tyr Arg 260 265 270
- Pro Ser Gly Ser Gly Met Thr Ser Leu Val Thr Leu Phe Leu Gln Gly 275 280 285
- Thr Gln Val Ser Gly Thr Leu Pro Pro Gln Trp Arg Ser Met Leu Asn 290 295 300
- Ala Glu Phe Leu Gln Leu Glu Asn Cys Asp Leu Ser Gly Cys Leu Pro 305 310 315
- Pro Glu Trp Ala Ala Met Pro Lys Leu Arg His Val Glu Leu Lys Gly 325 330 335
- Asn Gln Phe Ala Gly Cys Val Pro Asp Ser Trp Ala Gln Lys Ala Gly 340 345 350
- Leu Val Val Glu Ile Glu Asp Lys His Thr Gly Asn Ser Cys Ile Ala 355 360 365
- Gly Ala Asp Cys Ala Thr Thr Thr Thr Thr Thr Glu Pro Thr Ser 370 375 380
- Thr Ala Ser Pro Thr Ala Thr Pro Thr Ser Ala Pro Glu Thr Glu Cys 385 390 395
- Glu Val Asp Gly Cys Glu Val Cys Asp Gly Asp Ser Ala Ala Arg Cys
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Ala Arg Cys Arg Glu Gly Tyr Phe Leu Thr Asp Glu Arg Thr Cys Leu 420 425 430

Val Tyr Arg Asp Gly Gly Val Val Ala Val Ser Ile Gly Ala Ala Ala 435 440 445

Ala Ala Val Val Cys Met Ala Val Leu Leu Ser Val Gly Leu Ala Ala 450 455 460

